RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/539,440
Source: PCT
Date Processed by STIC: D13/06

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PCT

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PATENT APPLICATION: US/10/539,440
                                                            TIME: 09:28:18
                     Input Set : A:\SEQ list P697PC00.ST25.txt
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      5 <110> APPLICANT: ENKAM Pharmaceutical A/S
      9 <120> TITLE OF INVENTION: Method of modulation of interaction between receptor and
ligand
    13 <130> FILE REFERENCE: P 697 PC00
C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/539,440
C--> 17 <141> CURRENT FILING DATE: 2005-06-20
     17 <160> NUMBER OF SEQ ID NOS: 146
     21 <170> SOFTWARE: PatentIn version 3.1
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     29 <212> TYPE: PRT
     31 <213> ORGANISM: Artificial sequence
     35 <220> FEATURE:
     37 <223> OTHER INFORMATION: NCAM Fn III, 2 [Swiss-Prot: P13591]: FGFR binding motif
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     41 Glu Val Tyr Val Val Ala Glu Asn Gln Gln Gly Lys Ser Lys Ala
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              ing motif
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FGFR
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              binding motif
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     83 Ala Thr Asn Arg Gln Gly Lys Val Lys Ala Phe Ala His Leu
     84 1
                                            10
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     91 <212> TYPE: PRT
     93 <213> ORGANISM: Artificial sequence
     97 <220> FEATURE:
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99 <223> OTHER INFORMATION: Disintegrin and metalloprotease domain 8 (ADAM-8) [Swiss-

Prot: Q0

100 5910]: FGFR binding motif

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     112 <212> TYPE: PRT
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     118 <220> FEATURE:
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     121
               binding motif
     123 <400> SEQUENCE: 5
     125 Val Ala Glu Asn Ser Arg Gly Lys Asn Val Ala Lys Gly
                         5
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     133 <212> TYPE: PRT
     135 <213> ORGANISM: Artificial sequence
     139 <220> FEATURE:
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FGFR b
     142
               inding motif
     144 <400> SEQUENCE: 6
     146 Gly Glu Tyr Trp Cys Val Ala Glu Asn Gln Tyr Gly Gln Arg
     147 1
     150 <210> SEQ ID NO: 7
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     154 <212> TYPE: PRT
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     162 <223> OTHER INFORMATION: FIII,1 domain of NCAM [Swiss-Prot: P13591]: FGFR binding
motif
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     166 Arg Leu Ala Ala Leu Asn Gly Lys Gly Leu Gly Glu Ile Ser
     167 1
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     171 <210> SEQ ID NO: 8
     173 <211> LENGTH: 16
     175 <212> TYPE: PRT
     177 <213> ORGANISM: Artificial sequence
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     183 <223> OTHER INFORMATION: Neuronal nicotinic acetylcholine receptor alpha 3 subunit
(CHRNA
               3) [Swiss-Prot: Q8VHH6/P04757:/Q8R4G9/P32297]: FGFR binding motif
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     186 <400> SEQUENCE: 8
     188 Lys Tyr Ile Ala Glu Asn Met Lys Ala Gln Asn Val Ala Lys Glu Ile
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     194 <211> LENGTH: 14
     196 <212> TYPE: PRT
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motif
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206 <400> SEQUENCE: 9

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     218 <213> ORGANISM: Artificial sequence
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CSF-
     225
               R; CD114 antigen) [ Swiss-Prot: Q99062]: FGFR binding motif
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     229 Lys Gly Leu Gly Glu Ile Ser Ala Ala Thr Glu Phe Lys Thr
     230 1
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     239 <213> ORGANISM: Artificial sequence
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     278 <212> TYPE: PRT
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075462]: FG
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     291 Glu Ile Trp Val Glu Ala Thr Asn Arg Leu Gly
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     299 <212> TYPE: PRT
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     305 <220> FEATURE:
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     325 <220> FEATURE:
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Prot: O9D
     328
               CM6]: FGFR binding motif
     330 <400> SEQUENCE: 15
     332 Glu Val Trp Ile Glu Lys Asp Pro Ala Lys Gly Arg Ile
     336 <210> SEQ ID NO: 16
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     340 <212> TYPE: PRT
     342 <213> ORGANISM: Artificial sequence
     346 <220> FEATURE:
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binding
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     351 <400> SEQUENCE: 16
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FGFR
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     374 Lys Tyr Val Glu Leu Tyr Leu Val Ala Asp Tyr Leu Glu Phe Gln Lys
     375 1
                                              10
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binding
     391
               motif
     393 <400> SEQUENCE: 18
     395 Arg Tyr Val Glu Leu Tyr Val Val Val Asp Asn Ala Glu Phe Gln
     396 1
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     399 <210> SEO ID NO: 19
     401 <211> LENGTH: 16
     403 <212> TYPE: PRT
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     409 <220> FEATURE:
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414 <400> SEQUENCE: 19

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     426 <213> ORGANISM: Artificial sequence
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TECADAM [
               AF163291] : FGFR binding motif
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     435 <400> SEQUENCE: 20
     437 Lys Tyr Ile Glu Tyr Tyr Val Val Leu Asp Asn Gly Glu Phe Lys Lys
     438 1
                                              10
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     445 <212> TYPE: PRT
     447 <213> ORGANISM: Artificial sequence
     451 <220> FEATURE:
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FGFR
     454
               binding motif
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     458 Arg Tyr Leu Glu Leu Tyr Ile Val Ala Asp His Thr Leu Phe
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     466 <212> TYPE: PRT
     468 <213> ORGANISM: Artificial sequence
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     474 <223> OTHER INFORMATION: ADAM-1A Fertilin alpha [Swiss-Prot: Q8R533]: FGFR binding
motif
     476 <400> SEOUENCE: 22
     478 Lys Tyr Val Glu Met Phe Val Val Val Asn His Gln Arg Phe Gln
     479 1
                         5
                                              10
     482 <210> SEQ ID NO: 23
     484 <211> LENGTH: 14
     486 <212> TYPE: PRT
     488 <213> ORGANISM: Artificial sequence
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     499 1
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     504 <211> LENGTH: 16
     506 <212> TYPE: PRT
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     512 <220> FEATURE:
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     516 <400> SEQUENCE: 24
     518 Lys Tyr Val Glu Leu Phe Ile Val Ala Asp Asp Thr Val Tyr Arg Arg
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VERIFICATION SUMMARY

DATE: 07/13/2006 TIME: 09:28:19

PATENT APPLICATION: US/10/539,440

Input Set : A:\SEQ list P697PC00.ST25.txt Output Set: N:\CRF4\07132006\J539440.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application No L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date